

Serial Number: 10/031,598

CRF Processing Date: 1/26/2002
 Edited by: _____
 Verified by: [Signature] (STIC staff)

ENTERED

- ☐ Changed a file from non-ASCII to ASCII. ENTERED

☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically: # 2

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

¹Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

Inserted colons after headings/subheadings. Headings edited included:

Deleted extra, invalid, headings used by an applicant, specifically:

Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file;
☐ page numbers throughout text; ☐ other invalid text, such as _____.

Inserted mandatory headings, specifically:

Corrected an obvious error in the response, specifically:

Edited identifiers where upper case is used but lower case is required, or vice versa.

Corrected an error in the Number of Sequences field, specifically:

A "Hard Page_Break" code was inserted by the applicant. All occurrences had to be deleted.

Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

Other:

***Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.** 3/1/95

3/1/95



PCF/10

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/10/037,598

DATE: 01/26/2002
 TIME: 16:29:31

Input Set : A:\PTO.AMC.txt
 Output Set: N:\CRF3\01252002\J037598.raw

P.S

3 <110> APPLICANT: Monsanto Co
 4 Concibido, Vergel
 5 Delanney, Xavier
 7 <120> TITLE OF INVENTION: Soybean Plants with Enhanced Yields and Methods for Breeding
 for and
 8 Screening of Soybean Plants with Enhanced Yields
 10 <130> FILE REFERENCE: 38-21(52175)B
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/037,598
 C--> 12 <141> CURRENT FILING DATE: 2002-01-04
 12 <150> PRIOR APPLICATION NUMBER: 60/260,040
 13 <151> PRIOR FILING DATE: 2001-01-05
 15 <160> NUMBER OF SEQ ID NOS: 37
 17 <170> SOFTWARE: PatentIn version 3.0
 19 <210> SEQ ID NO: 1
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 21 <212> TYPE: DNA
 22 <213> ORGANISM: Glycine max
 24 <400> SEQUENCE: 1
 25 ggcgcgacaac tctaatagaaa atct 24
 28 <210> SEQ ID NO: 2
 29 <211> LENGTH: 23
 30 <212> TYPE: DNA
 31 <213> ORGANISM: Glycine max
 33 <400> SEQUENCE: 2
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 37 <210> SEQ ID NO: 3
 38 <211> LENGTH: 25
 39 <212> TYPE: DNA
 40 <213> ORGANISM: Glycine max
 42 <400> SEQUENCE: 3
 43 gcgttttaaat ttatgatata accaa 25
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 47 <211> LENGTH: 24
 48 <212> TYPE: DNA
 49 <213> ORGANISM: Glycine max
 51 <400> SEQUENCE: 4
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 55 <210> SEQ ID NO: 5
 56 <211> LENGTH: 25
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 58 <213> ORGANISM: Glycine max
 60 <400> SEQUENCE: 5
 61 atcaatcgac gcaataatca agaaa 25
 64 <210> SEQ ID NO: 6

RAW SEQUENCE LISTING

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73 <210> SEQ ID NO: 7
74 <211> LENGTH: 25
75 <212> TYPE: DNA
76 <213> ORGANISM: Glycine max
78 <400> SEQUENCE: 7
79 caggcttcag tgtgcataat acagg 25
82 <210> SEQ ID NO: 8
83 <211> LENGTH: 25
84 <212> TYPE: DNA
85 <213> ORGANISM: Glycine max
87 <400> SEQUENCE: 8
88 ttctatgttc cctgtgcaaa cactg 25
91 <210> SEQ ID NO: 9
92 <211> LENGTH: 25
93 <212> TYPE: DNA
94 <213> ORGANISM: Glycine max
96 <400> SEQUENCE: 9
97 gtctgcaagc taacagtgtc agagg 25
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101 <211> LENGTH: 26
102 <212> TYPE: DNA
103 <213> ORGANISM: Glycine max
105 <400> SEQUENCE: 10
106 cacactcaat ctcattagca gacacg 26
109 <210> SEQ ID NO: 11
110 <211> LENGTH: 25
111 <212> TYPE: DNA
112 <213> ORGANISM: Glycine max
114 <400> SEQUENCE: 11
115 tcctttggct cactattgac gattt 25
118 <210> SEQ ID NO: 12
119 <211> LENGTH: 25
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121 <213> ORGANISM: Glycine max
123 <400> SEQUENCE: 12
124 acccgtgtgc cactttaact acatt 25
127 <210> SEQ ID NO: 13
128 <211> LENGTH: 25
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132 <400> SEQUENCE: 13
133 taacgctgca tgatttgagt tctgt 25
136 <210> SEQ ID NO: 14
137 <211> LENGTH: 25

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146 <211> LENGTH: 28
147 <212> TYPE: DNA
148 <213> ORGANISM: Glycine max
150 <400> SEQUENCE: 15
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154 <210> SEQ ID NO: 16
155 <211> LENGTH: 28
156 <212> TYPE: DNA
157 <213> ORGANISM: Glycine max
159 <400> SEQUENCE: 16
160 gcgatgctta cttttcctat gatcactt 28
163 <210> SEQ ID NO: 17
164 <211> LENGTH: 24
165 <212> TYPE: DNA
166 <213> ORGANISM: Glycine max
168 <400> SEQUENCE: 17
169 gcgtagcaac aaagcaatct acag 24
172 <210> SEQ ID NO: 18
173 <211> LENGTH: 29
174 <212> TYPE: DNA
175 <213> ORGANISM: Glycine max
177 <400> SEQUENCE: 18
178 gcgtcccatt ttattccaca ctatgtaat 29
181 <210> SEQ ID NO: 19
182 <211> LENGTH: 235
183 <212> TYPE: DNA
184 <213> ORGANISM: Glycine max
186 <400> SEQUENCE: 19
187 cgacaactct aatgaaaatc tttattatta ttattattat tattattatt attattattc 60
189 acgaagttcc cttaaaaaat ctttagtaag acacatgcat taattatatg acaataaaaa 120
191 aaaaaagaat tcaaatgttt caaaatgaaa aatcattaat tcacttttat gtcaattatt 180
193 attattatta ttataacatt aattactttg aattgacttt tgaaaaatca aactc 235
196 <210> SEQ ID NO: 20
197 <211> LENGTH: 272
198 <212> TYPE: DNA
199 <213> ORGANISM: Glycine max
201 <400> SEQUENCE: 20
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204 attattatta ttattattat tattattaaa agttatacat gtaaatatth ttttaagggtg 120
206 acattctgaa taaattttta tatgtgattt gggaaaagta gagacaagtt caccctaaaa 180
208 ttaatattca gtaagtggaa cgtctccaaa ttattataaa aaattgtaaa tttttattct 240
210 atgcgactga agttgtggaa aaagagataa aa 272
213 <210> SEQ ID NO: 21
214 <211> LENGTH: 280

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221 attatatata tatatatata tatatatata tatatatata tatatatata gacaccccaa      120
223 taaaaatcat attaaaacaa ttataattca taatattcag aataaataaa aatattgaaa      180
225 taaatggcaa cacctcatcg tattcaaata aatataattg acacaacttt atactcaatt      240
227 ttttggttcc tggaatgaca tcccattgtc ttctcatcat      280
230 <210> SEQ ID NO: 22
231 <211> LENGTH: 366
232 <212> TYPE: DNA
233 <213> ORGANISM: Glycine max
235 <400> SEQUENCE: 22
236 caggcttcag tgtgcataat acagggtttct gttgggtggga ctttctccca acatttcatt      60
238 ttgggatttt ctcccaacct ttattttgtc tgaccttagt cgtaaatagt ctaaccttcc      120
240 ttcccttcctt catgtttcat tcgtgatcct gttttttggt atttcagggg gttggttgag      180
242 cctagtaggg ggccagggtg caacctatag ttgggatttc accccttagg ctgaaatttc      240
244 ctttctcac ttaagtaaaa aaaaaaacia aaagttttag tttttgtatg aaaatgcttt      300
246 tttatagcaa ttttatatga ttagaaaatt aaactattcc ccagtgtttg cacagggaac      360
248 atagaa      366
251 <210> SEQ ID NO: 23
252 <211> LENGTH: 96
253 <212> TYPE: DNA
254 <213> ORGANISM: Glycine max
256 <400> SEQUENCE: 23
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259 aatgatgaaa cgtgtctgct aatgagattg agtgtg      96
262 <210> SEQ ID NO: 24
263 <211> LENGTH: 321
264 <212> TYPE: DNA
265 <213> ORGANISM: Glycine max
267 <400> SEQUENCE: 24
268 tcctttggct cactattgac gattttctcg atgattaatt gacccaacat tctgtttgta      60
270 actttattta taaaacaaat atttgtactt caattataac aacaaattta agaagaatat      120
272 atatatatat atatttgtga tggaaatgat catgaaagaa acagaatcaa tatttcttat      180
274 aatcaagaaa aataatagac tcattttattt cttataaaaa gaaggagata aagtataaaa      240
276 tacaatggt aaacataaaa gaaaaaaaaa ctttttttga ccggtatggt aacgaaaatg      300
278 tagttaaagt ggcacacggg t      321
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283 <212> TYPE: DNA
284 <213> ORGANISM: Glycine max
286 <400> SEQUENCE: 25
287 taacgctgca tgatttgagt tctgttttgt cggcggggac tagggacaaa tatatttttt      60
289 gttagttaat ttgtatatatt attggtgata tgtctgaagt taagttaatt ggccatgcat      120
291 gtgtgtgtgt gtggtagtga gaagaattga gaaaaagaat gtggtctcca aagtccaacc      180
293 aatac      185
296 <210> SEQ ID NO: 26
297 <211> LENGTH: 3830

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298 <212> TYPE: DNA
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304 aatgtcaagt gagtttagaa tactaaatga aaattttaac ataaaaaaaa aaaaatcaat 120
306 ggaatggaac ccatccagcg caactagctg agtcacatac agtgccaaaa gacatgggta 180
308 ctacaaatgc tcactttagt ggctatggaa caaccatcag cattcagctc ttccotTTTT 240
310 ctgtcgtagg ccaagagaca aagtttgtca caggtttaca aattgattgt ggccacaatc 300
312 acacggtaaa cattagaatg gaagaaaaaa aatctgtcta tgatcgatgt cgtgaacttc 360
314 acccactcca tcaatgaaga atttatttta aatacagtta cacaccaact taataagact 420
316 ttttgcacaa aattacctga ttgggaggaa tatgaattgt cttataaatc acgtattcac 480
318 aagttctact tttaaaaaac totttacatg tattttccaa aaaaagaaaa atctttacat 540
320 gtatgttaac ctacctaaca aatctctaata taacctataa attttttaaa tgctttttga 600
322 gaaaacttta taggcagata gaagattggt gagagtTTTT taaatgctta tcaacaatct 660
324 cogatagtcc cttagcttta ccaagtacat gaaaatctta catataatgc ttttacttta 720
326 ccaactatta acttgagcac cgaaatcttt accagtatgc tcatttgatg catattaaaa 780
328 tgtacaaaat tttatagagg cctgatcaat accatcgaat gaaaccttaa tgacatgcta 840
330 cttgttagcg atgtcaataa aggttacttc aaggattatt ccacaggcct aaatcataga 900
332 caattttact taattgtatt tattcaatta gtccttagat gtcaaagaat ctattagatg 960
334 atagttttag tggcatgata gagaatgaaa cccacatcta taaaaaaaag aagacaaaag 1020
336 ttagttttag atctttaatc acttggtgta attcatatta gttttacgtg tattcgaagt 1080
338 gaaaatatct atctgtatga gaccataaac attcctatga gagacttggt tgaagtataa 1140
340 tttttcatag tacagtaaaag ctgattggtg ttttttctcg tacgcaaaat ttatattcag 1200
342 gacaatgttt aagagtgaag acataataaa attaacctca caaaaagtaa gtatatatat 1260
344 atatatatat atatatatat atataataat ctcaatcaat taaaataata ataaggacaa 1320
346 ataaatagat tctcacaaaa tataatttat tattaaatta atttttaaca ttataactta 1380
348 acgataaaat atttttttta ttttttttta tgaactaatt taacaactca tcacatcttg 1440
350 caaaacaaaa tgaatcattt atcctaataa taatttaatt taggcgttta ttttatgatg 1500
352 atttagcatc tttttgggag aatactaaaa aacatataaa agaaaaagaa atattcagga 1560
354 tgaaaaatga aatgcgtgtg aaaattggaa ggaggtaagg ctgggtcgac ccagatctag 1620
356 ttgagctcac caactcccg cccattttcc ttatttatag acagagtctg attgtttcct 1680
358 caccactccc tccactctct ttctctagtc ctgttatttc tcagcgcgta aagcatggct 1740
360 ttgttggtgg agaaaaccac gagtggtcgc gagtacaagg tcaaggacct ttccaggcc 1800
362 gacttcggcc gcctcgagat cgagctggcc gaggttgaga tgcccgccct catggcctgt 1860
364 cggaccgagt tcggcccttc ccagcccttc aagggggccc gcatcacccg ctccctccac 1920
366 atgaccatcc agaccgccgt tctcattgag accctcaccg cccttgccgc cgaggtccgc 1980
368 tgggtgctct gcaacatctt ctccacccag gaccacgccg ccgcccgtat tgcccgcgac 2040
370 agtgccgcgc tcttcgcctg gaagggtgag accctccagg agtactggtg gtgcaccgag 2100
372 cgcgccctcg actggggccc cgggtggtgga cccgacctca tcgtcgacga cgggtggtgac 2160
374 gctacccttc tcatccacga aggcgtcaag gccgaggagc tctatgagaa gaccggcgaa 2220
376 ctccccgacc ccaactccac cgacaacgcc gagtttcaga tcgtgcttac catcatcaga 2280
378 gatgggttga agaccgatcc caccaggtag cgcaagatga aggagcgtct cgttgggggt 2340
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382 ctcttccctg ctattaatgt caatgactct gtcaccaaga gcaaggtaat gtctcttttt 2460
384 cccccagatc tagtgtcttt tttgtgttaa aatgtaggat tgagttcgga tctgttgttt 2520
386 ttggatgggt tttgtgccat tggtgaaatg aggttttgaa cctgtcaact gtttgactaa 2580
388 tgtcctctaa gaagtctgga tcggtatttg gtgctatttt agtgtgtttg gatctgtgtg 2640
390 ttgaaacgtc agaacttag taagttgctt gctaacgtga ctttaggtaa atggtcacat 2700
392 gttttattac acaaataagg aattgattct gagtgcacat tttgatttga agctactttt 2760

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→ Use of n and/or Xaa has been detected in the Sequence Listing.
 Review the Sequence Listing to insure a corresponding
 explanation is presented in the <220> to <223> fields of
 each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/037,598

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Input Set : A:\PTO.AMC.txt

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L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:1336 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34

L:1338 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34

L:1340 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34

L:1346 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34